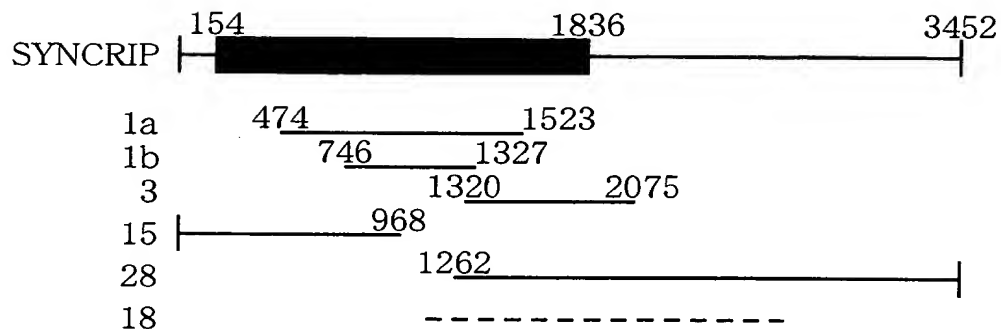


FIG.1

A



B

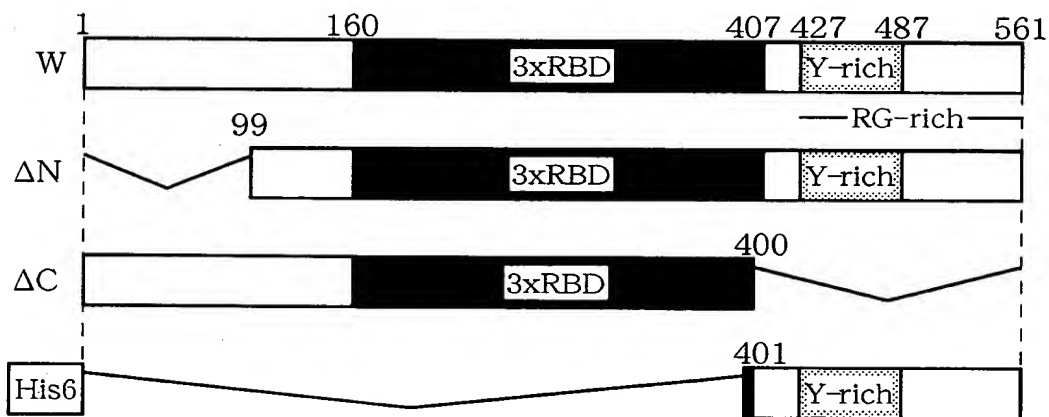


FIG.2

SYNCRIP 1: ---MATEHVNGNGTTEPMDDTSAVIESENFOITLDAGLEPKVAEKLDHTYVAGLVAHSD 56  
hnRNPR 1: MANOVNGNAVQLKEEPEPMD-TSSVTHTEHYKTLIEAGLEPKVAERLDEIFOTGLVAVVD 59

SYNCRIP 57: LDERATEALKEFNEDGALAVLQQPKDSDLSHVONKSAFLCGVMKTYRQREKQGTQVADSS 116  
hnRNPR 60: LDERAIDALRENEEGALSVLQPKESDLSHVONKSAFLCGVMKTYRQREKQGSKVQEST 119

SYNCRIP 117: KGPDEAKIKALLERTGYTLDVTTGQRKYGGPEPDSVYSGQOPSVGTEIFNGKLEPDLFED 176  
hnRNPR 120: KGPDEAKIKALLERTGYTLDVTTGQRKYGGPEPDSVYSGVOPGIGTEVEVGKLEPDLFED 179

SYNCRIP 177: ELVPLFEKAGPIWDLRLMMDPI/TGLNRYAFVTFCTKEAAQBAVKLYNNHRIKSGKHIGV 236  
hnRNPR 180: ELVPLFEKAGPIWDLRLMMDPI/SQNRGYAFITFCGKEAAQBAVKLCDSYIRPGKHIGV 239

SYNCRIP 237: CISVANNRLFVGSIFKSKTKKQILEEFSKVTEGLTDVILYHQPDDKKKNGGFCFLEYEDH 296  
hnRNPR 240: CISVANNRLFVGSIFKNKTENILEEFSKVTEGLVDVILYHQPDDKKKNGGFCFLEYEDH 299  
clone-18 1: -----DH 2  
(m-hnRNPR)

SYNCRIP 297: KTAQAARRRLMSGKVKVGNGVGVTEWADPIEDPDEVMKVKVLEVRNLATVTEELIK 356  
hnRNPR 300: KSAQAARRRLMSGKVKVGNGVVTEWADPVEEPPDEVMKVKVLEVRNLATVTEELIK 359  
clone-18 3: KSAQAARRRLMSGKVKVGNGVVTEWADPVEEPPDEVMKVKVLEVRNLATVTEELIK 62  
(m-hnRNPR)

SYNCRIP 357: SESEFGKLERVKKLDYAFVHEDRGAAVKAMDEMNGKEIEGEEIEVLAKPPDKKKRER 416  
hnRNPR 360: SESEFGKLERVKKLDYAFVHEDRGAAVKAMDEMNGKEIEGEEIEVLAKPPDKKKRER 419  
clone-18 63: SESEFGKLERVKKLDYAFVHEDRGAAVKAMDEMNGKEIEGEEIEVLAKPPDKKKRER 122  
(m-hnRNPR)

SYNCRIP 417: KAQRQAANKQMYDDYYHYGPPHMPPTRGGR-GGRGGYGYPPDYGYEDDY-DYGYDY 474  
hnRNPR 420: QAARQASRSTAYEDYYHYGPPHMPPIRGGRGGGRGGYGYPPDYGYEDDYDDYYGYDY 479  
clone-18 123: QAARQASRSTAYEDYYHYGPPHMPPIRGGRGGGRGGYGYPPDYGYEDDYDDYYGYDY 181  
(m-hnRNPR)

SYNCRIP 475: HNYRGGYEDPYGYED-FQVGAGRGGRGARGA-AFSRGRGAAPPRGRAGYSORGAPLGP 531  
hnRNPR 480: HDYRGGYEDPYGYDDGYAVRGRG-GGRGGRGAPPPRGRGAPPRGRAGYSORGAPLGP 538  
clone-18 182: HDYRGGYEDPYGYDDGYAVRGRG-GGRGGRGAPPPRGRGAPPRGRAGYSORGAPLGP 240  
(m-hnRNPR)

SYNCRIP 532: ARGVRGARGG-AQQQGRGGKGV-FA-GPDLLQ----- 561  
hnRNPR 539: PRGSRGGGGPAQQQGRGSRGSRGNRGGNVGGKRRKADGYNQPDSSKRRQTNNQONWGSQP 598  
clone-18 241: PRGSRGGGGPAQQQGRGSRGARGNRGNVGGKRRKADGYNQPDSSKRRQTNNQONWGSQP 300  
(m-hnRNPR)

562: -----  
hnRNPR 599: IAQQPLQGGDYSGNYGYNNDNQEFYQDTYGQQWK 633  
clone-18 301: IAQQPLQGGDYSGNYGYNNDNQEFYQDTYGQQWK 335  
(m-hnRNPR)

FIG.3

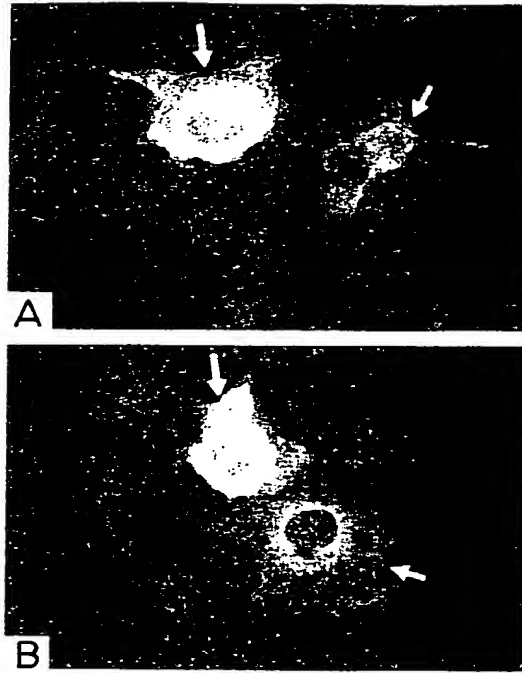


FIG.4

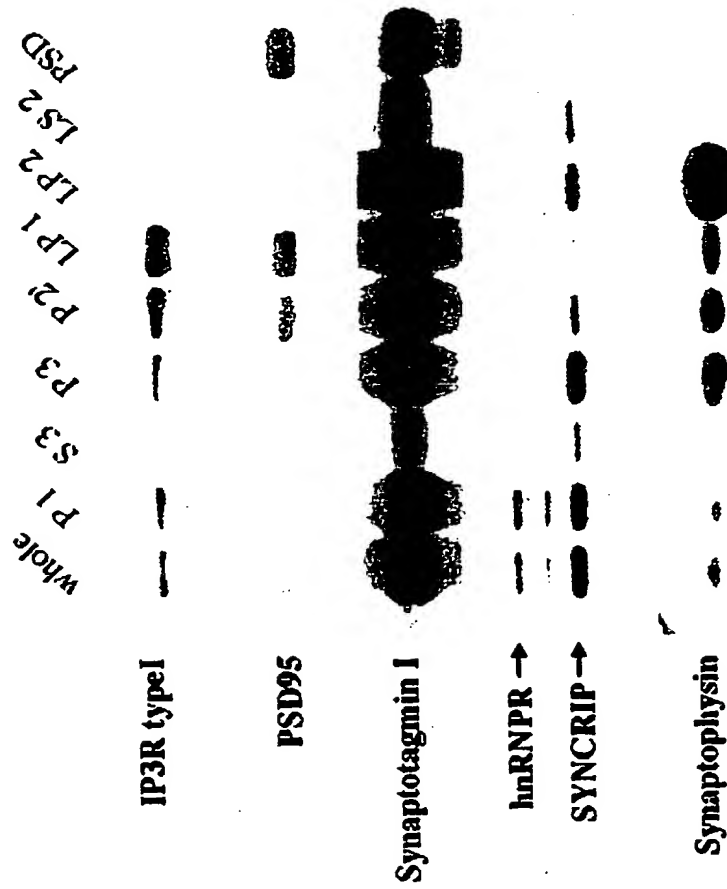


FIG.5

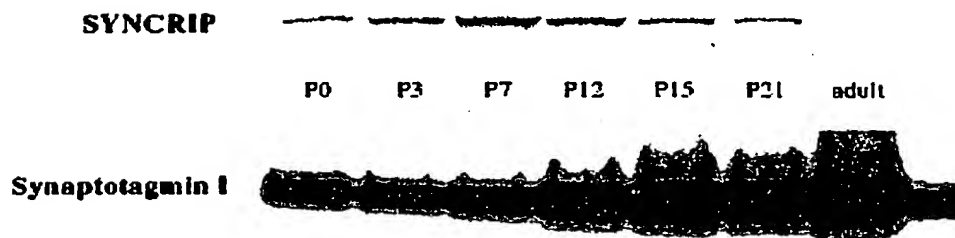


FIG.6

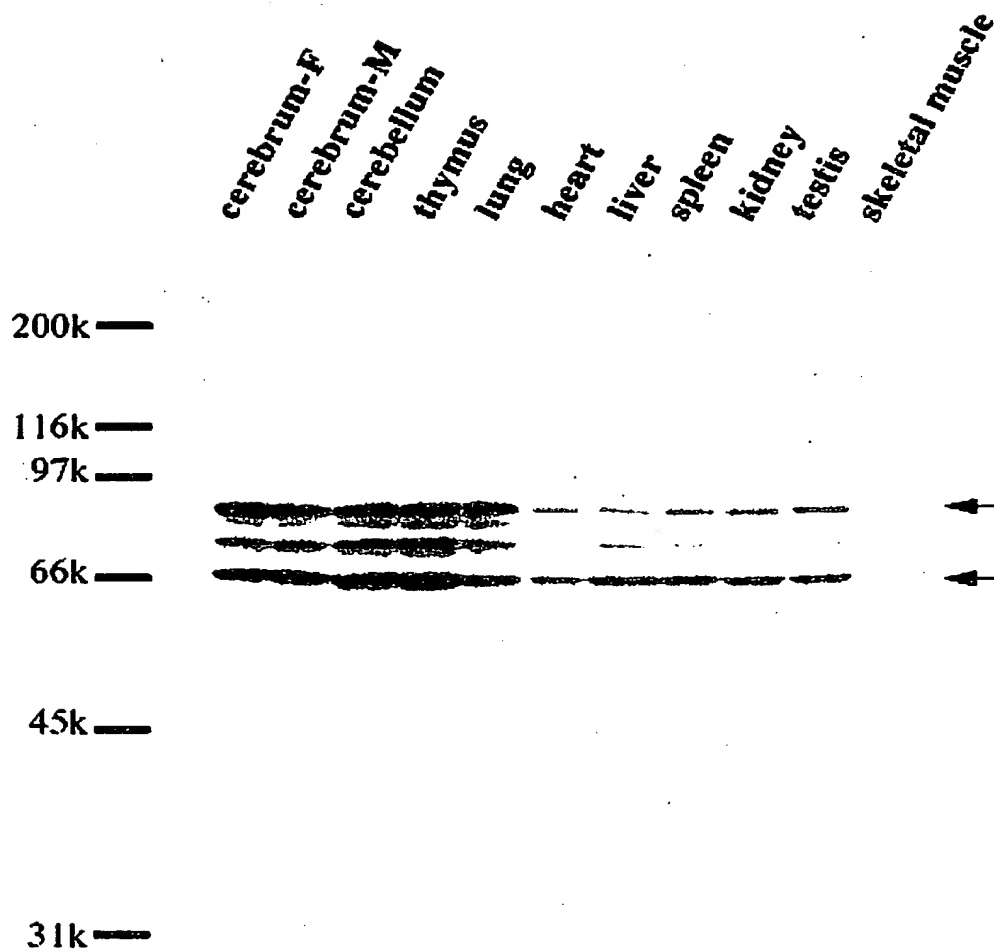


FIG.7

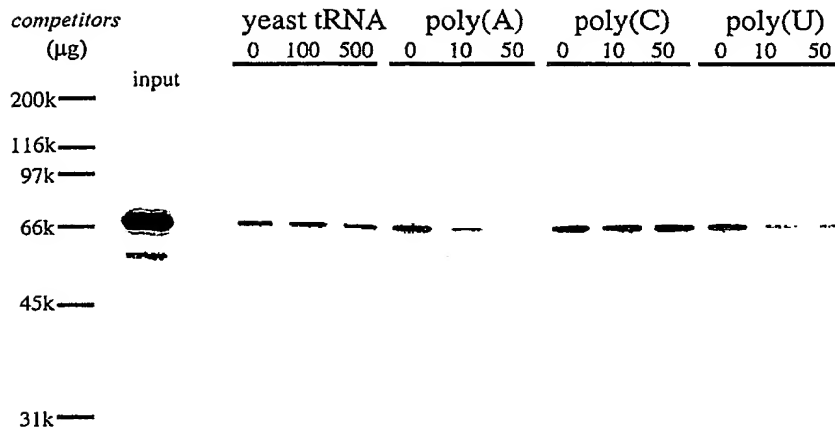
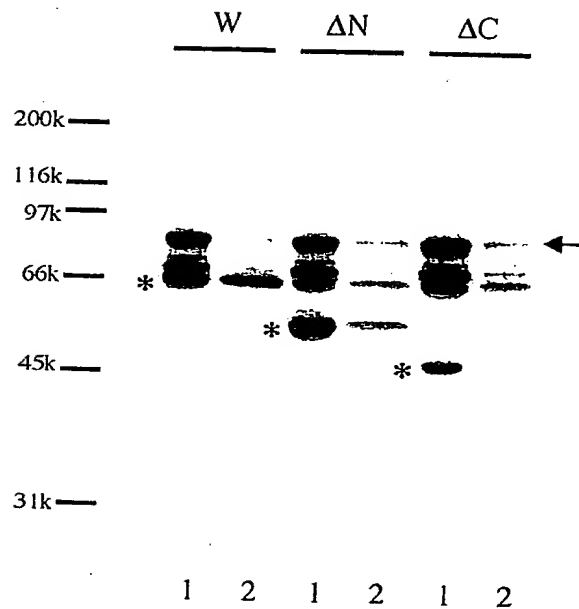


FIG.8





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FIG.9

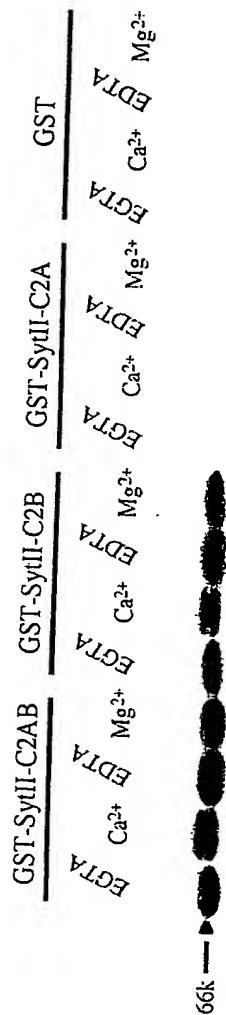


FIG.10

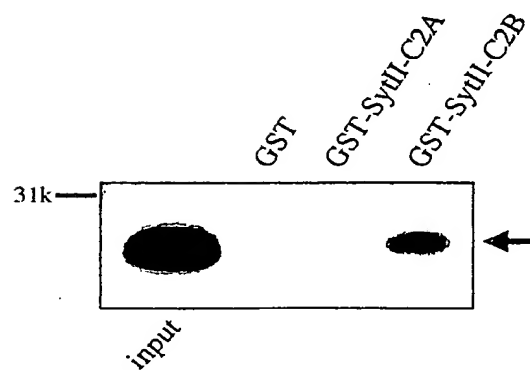


FIG.11

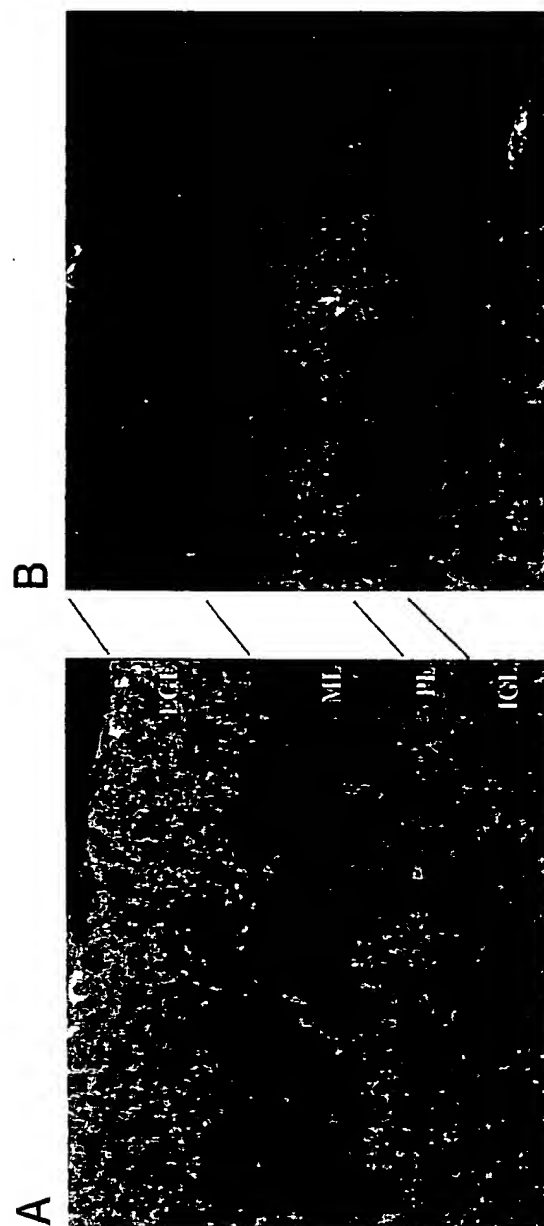


FIG.12

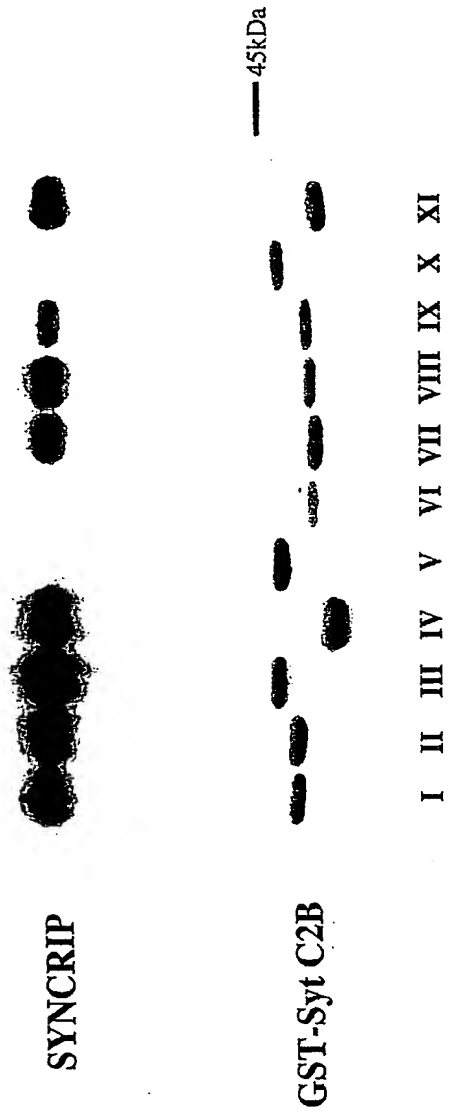


FIG.13

